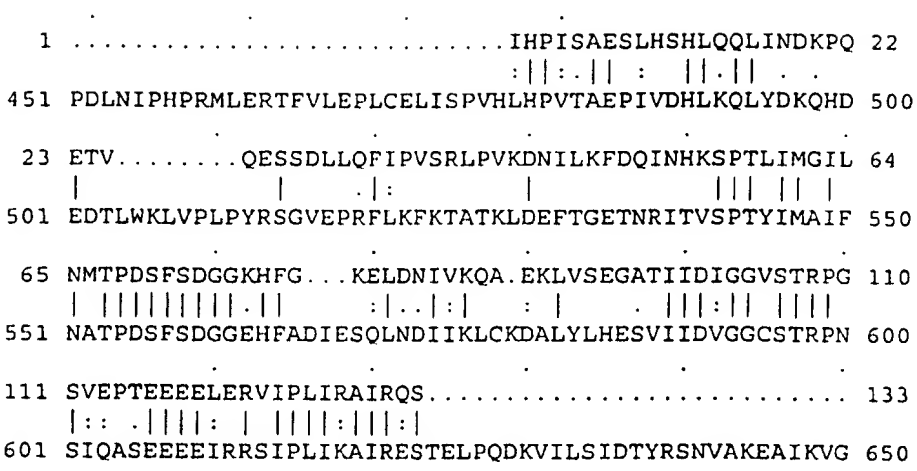
Fig.1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SEQIDNO:68
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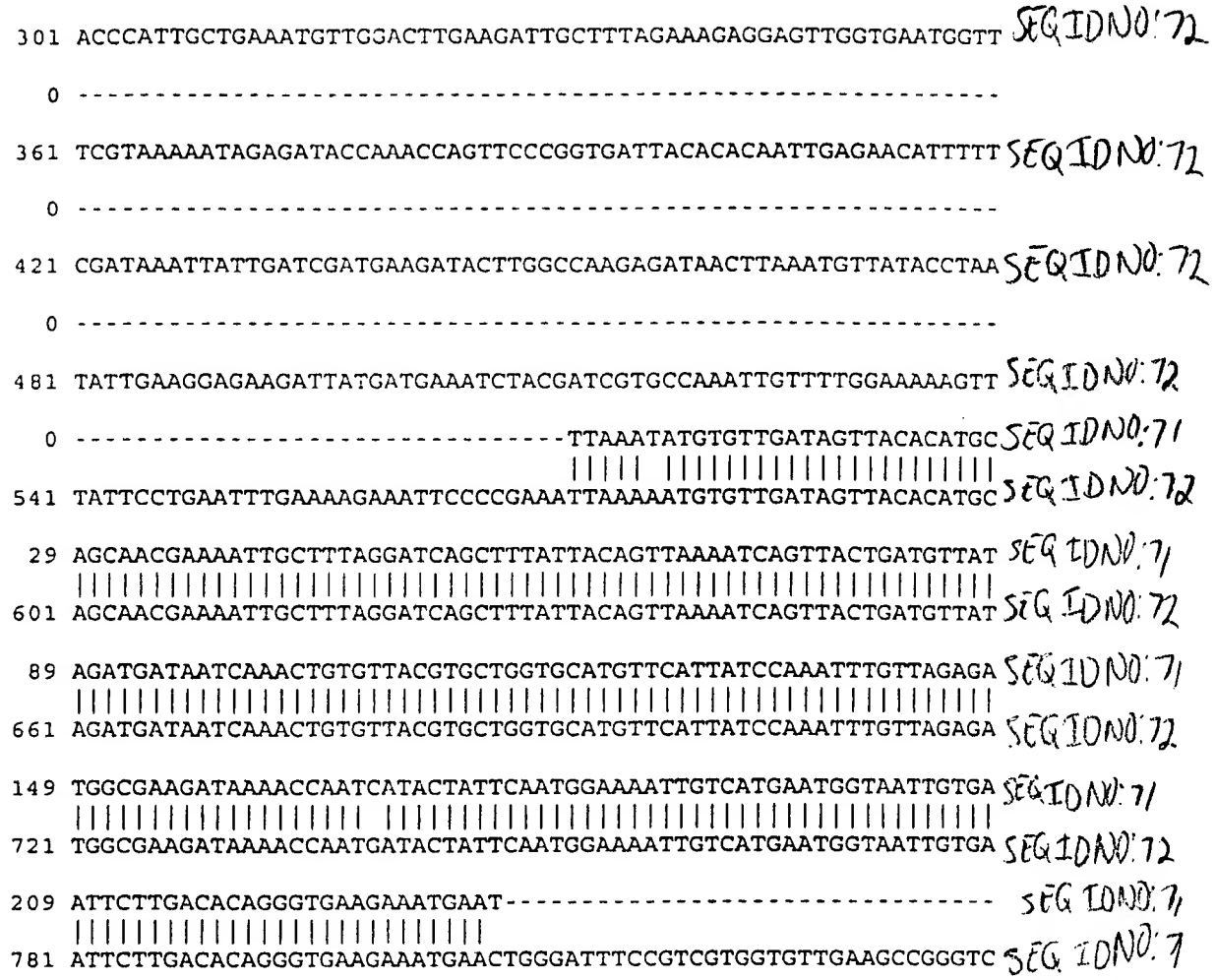
501 EDTLWKLVPLPYRSGVEPRFLKFKTATKLDEFTGETNRITVSPTYIMAIF 550
65 NMTPDSFSDGGKHFG...KELDNIVKQA.EKLVSEGATIIDIGGVSTRPG 110
551 NATPDSFSDGGEHFADIESQLNDIIKLCKDALYLHESVIIDVGGCSTRPN 600
111 SVEPTEEEELERVIPLIRAIRQS..... 133
601 SIQASEEEEIRRSIPLIKAIRESTELPQDKVILSIDTYRSNVAKEAIKVG 650

Fig.2

SEQ ID NO: 69

SEQ ID NO: 70

FIG.3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0 ----------
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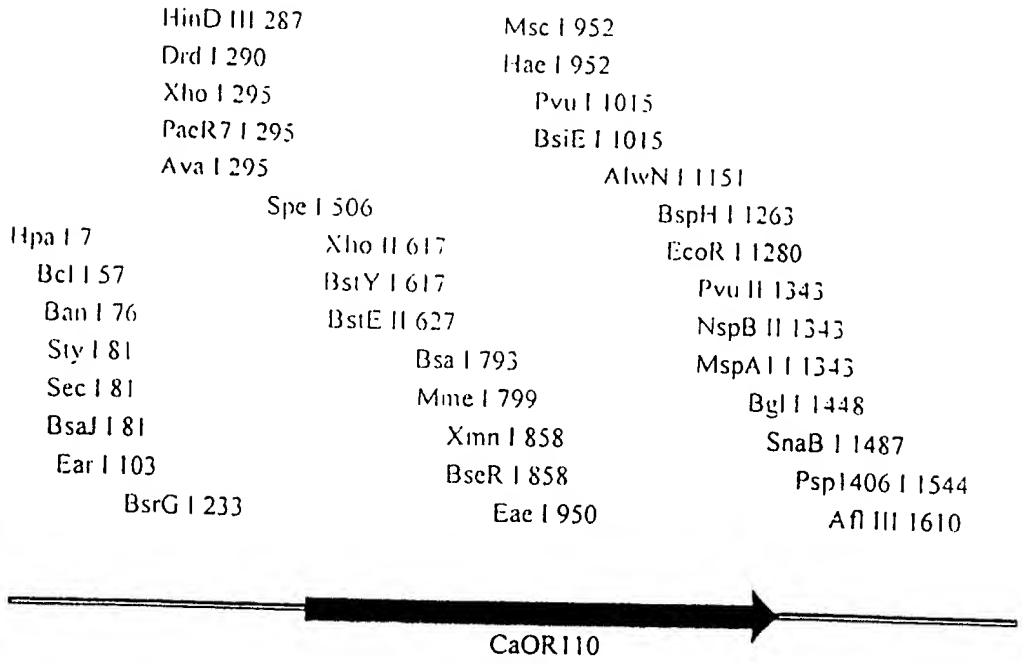
481 TATTGAAGGAGAAGATTATGATGAAATCTACGATCGTGCCAAATTGTTTTGGAAAAAGTT SEQ ID NO: 72
0 ----------TTAAATATGTGTTGATAGTTACACATGC SEQ ID NO: 71
541 TATTCCTGAATTTGAAAAGAAATTCCCCGAAATTAAAAATGTGTTGATAGTTACACATGC SEQ ID NO: 72
29 AGCAACGAAAATTGCTTTAGGATCAGCTTTATTACAGTTAAAATCAGTTACTGATGTTAT SEQ ID NO: 71
601 AGCAACGAAAATTGCTTTAGGATCAGCTTTATTACAGTTAAAATCAGTTACTGATGTTAT SEQ ID NO: 72
89 AGATGATAATCAAACTGTGTTACGTGCTGGTGCATGTTCATTATCCAAATTTGTTAGAGA SEQ ID NO: 71
661 AGATGATAATCAAACTGTGTTACGTGCTGGTGCATGTTCATTATCCAAATTTGTTAGAGA SEQ ID NO: 72
149 TGGCGAAGATAAAACCAATCATACTATTCAATGGAAAATTGTCATGAATGGTAATTGTGA SEQ ID NO: 71
721 TGGCGAAGATAAAACCAATGATACTATTCAATGGAAAATTGTCATGAATGGTAATTGTGA SEQ ID NO: 72
209 ATTCTTGACACAGGGTGAAGAAATGAAT---------- SEQ ID NO: 71
781 ATTCTTGACACAGGGTGAAGAAATGAACTGGGATTTCCGTCGTGGTGTTGAAGCCGGGTC SEQ ID NO: 7

FIG.4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PaeR7 I 295
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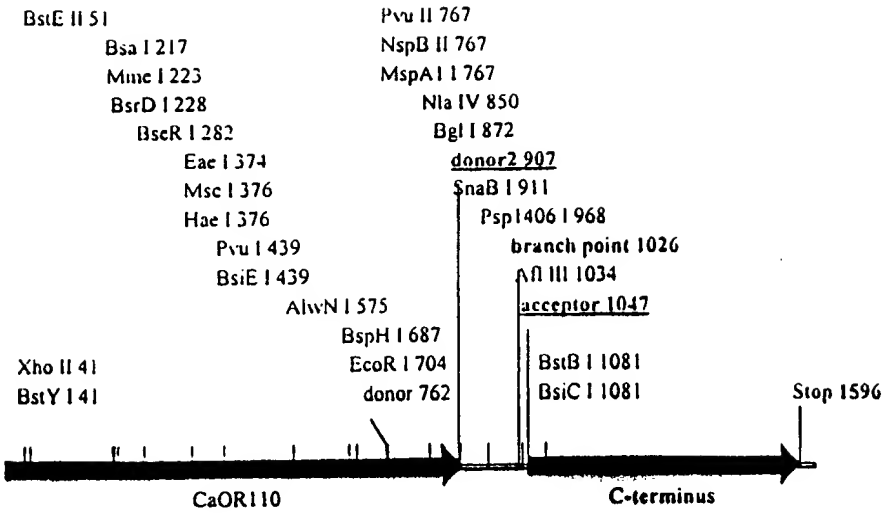
Sty I 81
Bsa I 793
Sec I 81
Mme I 799
BsaJ I 81
Xmn I 858
Ear I 103
BseR I 858
BsrG I 233
Eae I 950
Msc I 952
Hae I 952
Pvu I 1015
BsiE I 1015
AlwN I 1151
BspH I 1263
EcoR I 1280
Pvu II 1343
NspB II 1343
MspA I I 1343
Bgl I 1448
SnaB I 1487
Psp1406 I 1544
Afl III 1610
CaOR110

FIG.5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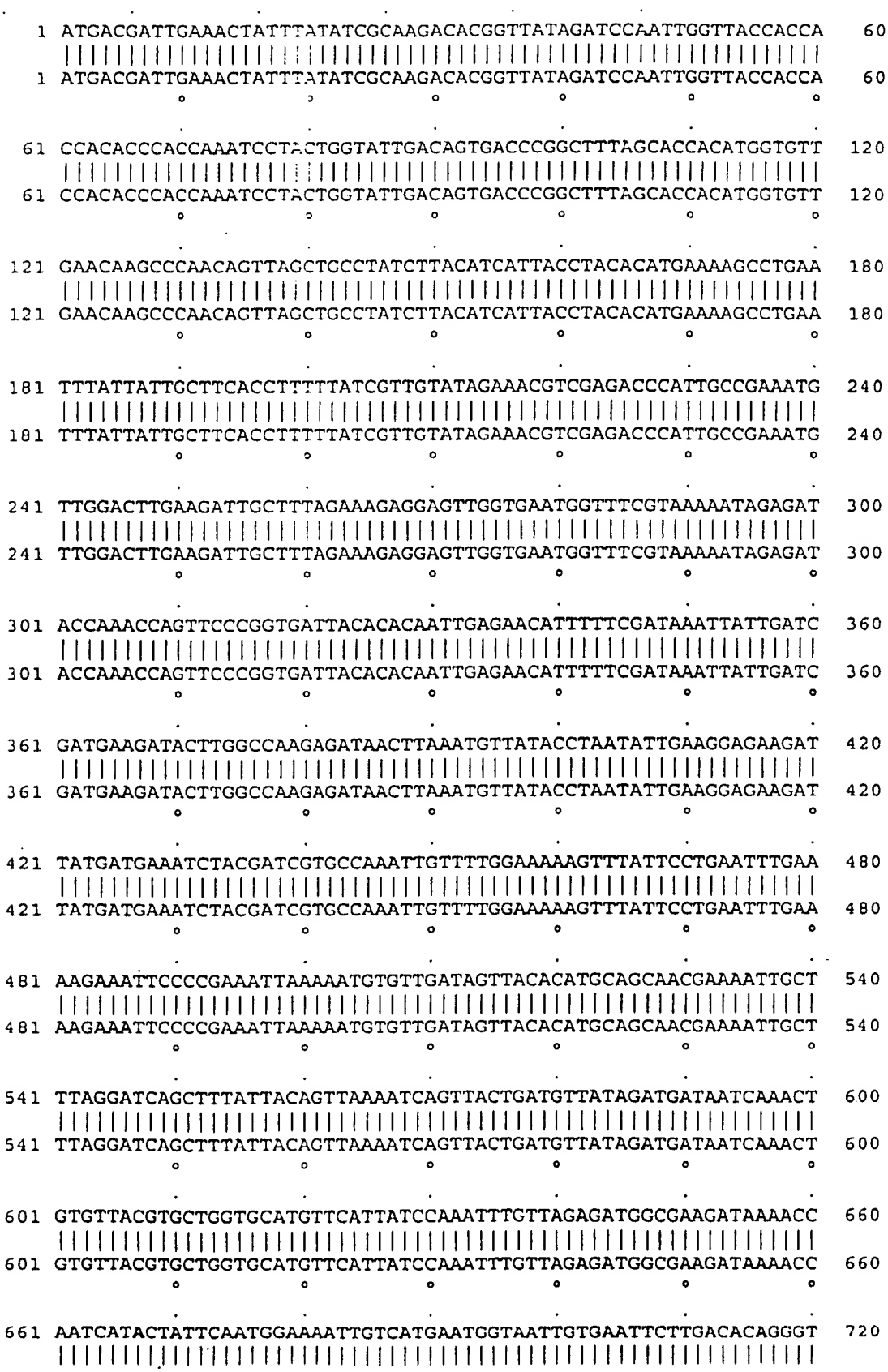
BspH I 687
EcoR I 704
donor 762
Xho II 41
BstY I 41
Pvu II 767
NspB II 767
MspA1 I 767
Nla IV 850
Bgl I 872
donor2 907
SnaB I 911
Psp1406 I 968
branch point 1026
Afl III 1034
acceptor 1047
BstB I 1081
BsiC I 1081
Stop 1596
CaOR110
C-terminus

FIG. 6A



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721 GAAGAAATGAACTGGGATTTCCGTCGTGGTGTTGAAGCCGGGTCAGCTGAAGATATAGCG  780

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    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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841 TCCGATGGTCCCATCACTGAATCTGCCACTGGGGCAGAAATAGATGGGAATGAAGATGAA  900
    ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
841 TCCGATGGTCCCATCACTGAATCTGCCACTGGGGCAGAAATAGATGGGAATGAAGATGAA  900

901 TTTGAAGTACGTAAAACTTGAAAGAGATATTAAATAGACACAAACTTAGAAAATATAGAG  960
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901 TTTGAA----------------------------------------------------  906

961 ATACAAACGTTTTGAATTTCTTGATTCACTTTTTTGTTTAAAAATAAAAATAGTTCAAAA 1020

906 ----------------------------------------------------------  905

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1081 CGAATAAAATCGACAATGAAGAAGAACCACCATCAAGGACAGGTCAAGCTCCAAAATTCA 1140
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1501 CTATCATGGATATAGATCAAGACTCACAAGGACAACAACCAGCTAGAAGTCAGTTCTTAA 1560
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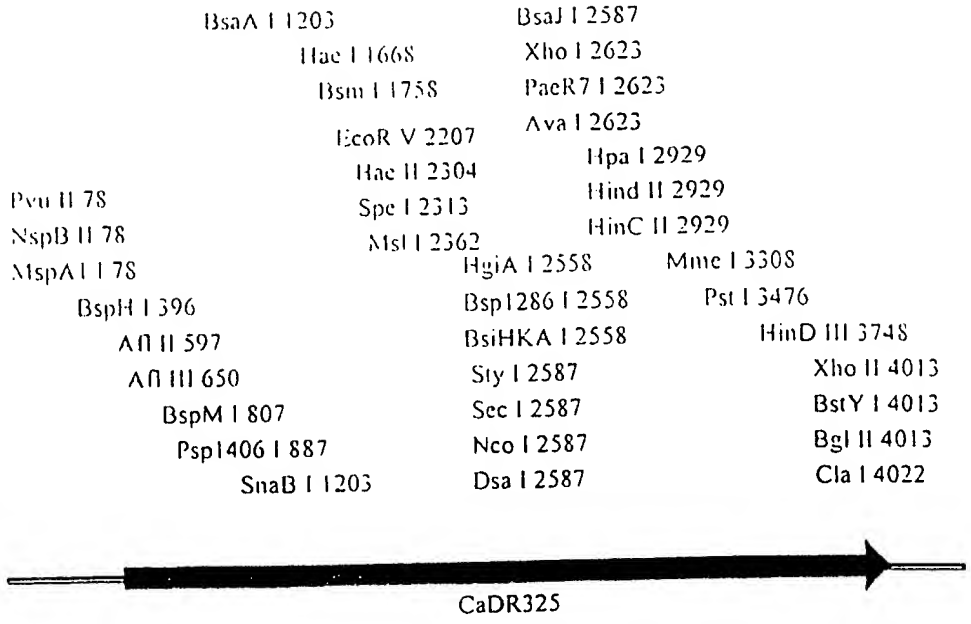
FIG. 7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Xho I 2623
PaeR7 I 2623
Ava I 2623
Hpa I 2929
Hind II 2929
HinC II 2929
Pvu II 78
NspB II 78
MspA1 I 78
BspH I 396
Afl II 597
Afl III 650
BspM I 807
Psp1406 I 887
SnaB I 1203
HgiA I 2558
Bsp1286 I 2558
BsiHKA I 2558
Sty I 2587
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Nco I 2587
Dsa I 2587
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